RAW SEQUENCE LISTING

The Biotechnology Systems Branch of the Scientific and Technical Information Center (STIC) no errors detected.

Application Serial Number: 10/031,158 B
Source: IFW0
Date Processed by STIC: 11/21/2005

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RAW SEQUENCE LISTING DATE: 11/21/2005
PATENT APPLICATION: **US/10/031,158B** TIME: 09:58:29

Input Set : **A:\sequence listing.txt**
Output Set: **N:\CRF4\11212005\J031158B.raw**

<110> APPLICANT: Pastan, Ira
Essand, Magnus
Lee, Byungkook
Vasmatzis, George
Wolfgang, Curt
The Government of the United States of America
as represented by the Secretary of the
Department of Health and Human Services
<120> TITLE OF INVENTION: T-Cell Receptor Gamma Alternate Reading Frame Protein,
(TARP) and Uses Thereof
<130> FILE REFERENCE: 4239-61854-01
<140> CURRENT APPLICATION NUMBER: 10/031,158B
<141> CURRENT FILING DATE: 2002-01-11
<150> PRIOR APPLICATION NUMBER: PCT/US00/19039
<151> PRIOR FILING DATE: 2000-07-12
<150> PRIOR APPLICATION NUMBER: US 60/157,471
<151> PRIOR FILING DATE: 1999-10-01
<150> PRIOR APPLICATION NUMBER: US 60/143,560
<151> PRIOR FILING DATE: 1999-07-13
<160> NUMBER OF SEQ ID NOS: 33
<170> SOFTWARE: PatentIn Ver. 2.1
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<212> TYPE: DNA
<213> ORGANISM: Artificial Sequence
<220> FEATURE:
<223> OTHER INFORMATION: Description of Artificial Sequence:PCR primer
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<213> ORGANISM: Artificial Sequence
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66 ttggacttgg attatcaaaa gtgg 24
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74 <220> FEATURE:
75 <223> OTHER INFORMATION: Description of Artificial Sequence:PCR primer
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78 ttgggcagtt ggaacaacct gaaa 24
81 <210> SEQ ID NO: 5
82 <211> LENGTH: 28
83 <212> TYPE: DNA
84 <213> ORGANISM: Artificial Sequence
86 <220> FEATURE:
87 <223> OTHER INFORMATION: Description of Artificial Sequence:PCR primer
89 <400> SEQUENCE: 5
90 gataaacaac ttgatgcaga tgtttccc 28
93 <210> SEQ ID NO: 6
94 <211> LENGTH: 28
95 <212> TYPE: DNA
96 <213> ORGANISM: Artificial Sequence
98 <220> FEATURE:
99 <223> OTHER INFORMATION: Description of Artificial Sequence:PCR primer
101 <400> SEQUENCE: 6
102 gggaaacatc tgcatcaagt tgtttatc 28
105 <210> SEQ ID NO: 7
106 <211> LENGTH: 27
107 <212> TYPE: DNA
108 <213> ORGANISM: Artificial Sequence
110 <220> FEATURE:
111 <223> OTHER INFORMATION: Description of Artificial Sequence:PCR primer
113 <400> SEQUENCE: 7
114 ctggagcttt gtttcagcaa ttgaagg 27
117 <210> SEQ ID NO: 8
118 <211> LENGTH: 27
119 <212> TYPE: DNA
120 <213> ORGANISM: Artificial Sequence
122 <220> FEATURE:
123 <223> OTHER INFORMATION: Description of Artificial Sequence:PCR primer
125 <400> SEQUENCE: 8
126 ctcaagaaga caaaggtatg ttccagc 27
129 <210> SEQ ID NO: 9
130 <211> LENGTH: 25
131 <212> TYPE: DNA
132 <213> ORGANISM: Artificial Sequence
134 <220> FEATURE:
135 <223> OTHER INFORMATION: Description of Artificial Sequence:PCR primer
137 <400> SEQUENCE: 9

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138 ttatgatttc tctccattgc agcag                                          25
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144 <213> ORGANISM: Artificial Sequence
146 <220> FEATURE:
147 <223> OTHER INFORMATION: Description of Artificial Sequence:PCR primer
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153 <210> SEQ ID NO: 11
154 <211> LENGTH: 24
155 <212> TYPE: DNA
156 <213> ORGANISM: Artificial Sequence
158 <220> FEATURE:
159 <223> OTHER INFORMATION: Description of Artificial Sequence:PCR primer
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165 <210> SEQ ID NO: 12
166 <211> LENGTH: 24
167 <212> TYPE: DNA
168 <213> ORGANISM: Artificial Sequence
170 <220> FEATURE:
171 <223> OTHER INFORMATION: Description of Artificial Sequence:PCR primer
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177 <210> SEQ ID NO: 13
178 <211> LENGTH: 1027
179 <212> TYPE: DNA
180 <213> ORGANISM: Homo sapiens
182 <220> FEATURE:
183 <221> NAME/KEY: CDS
184 <222> LOCATION: (74)..(247)
185 <223> OTHER INFORMATION: Coding region for PS-TCR gamma 1 polypeptide
186       (TARP)
188 <220> FEATURE:
189 <221> NAME/KEY: CDS
190 <222> LOCATION: (247)..(579)
191 <223> OTHER INFORMATION: Coding region for PS-TCR gamma 2 polypeptide (deduced amino
192       acid sequence not displayed along with DNA sequence, due to
193       overlapping CDS's)
196 <400> SEQUENCE: 13
197 gggcaagagt tgggcaaaaa aatcaaggta tttggtcccg gaacaaagct tatcattaca 60
199 gataaacaac ttg atg cag atg ttt ccc cca agc cca cta ttt ttc ttc    109
200                 Met Gln Met Phe Pro Pro Ser Pro Leu Phe Phe Phe
201                   1               5                  10
203 ctt caa ttg ctg aaa caa agc tcc aga agg ctg gaa cat acc ttt gtc   157
204 Leu Gln Leu Leu Lys Gln Ser Ser Arg Arg Leu Glu His Thr Phe Val
205          15                  20                  25
207 ttc ttg aga aat ttt tcc ctg atg tta tta aga tac att ggc aag aaa   205
```

```
208 Phe Leu Arg Asn Phe Ser Leu Met Leu Leu Arg Tyr Ile Gly Lys Lys
209      30                  35                  40
211 aga aga gca aca cga ttc tgg gat ccc agg agg gga aca cca           247
212 Arg Arg Ala Thr Arg Phe Trp Asp Pro Arg Arg Gly Thr Pro
213  45                  50                  55
215 tgaagactaa cgacacatac atgaaattta gctggttaac ggtgccagaa aagtcactgg 307
217 acaaagaaca cagatgtatc gtcagacatg agaataataa aaacggagtt gatcaagaaa 367
219 ttatctttcc tccaataaag acggatgtca tcacaatgga tcccaaagac aattgttcaa 427
221 aagatgcaaa tgatacacta ctgctgcagc tcacaaacac ctctgcatat tacatgtacc 487
223 tcctcctgct cctcaagagt gtggtctatt ttgccatcat cacctgctgt ctgcttagaa 547
225 gaacggcttt ctgctgcaat ggagagaaat cataacagac ggtggcacaa ggaggccatc 607
227 ttttcctcat cggttattgt ccctagaagc gtcttctgag gatctagttg ggctttcttt 667
229 ctgggtttgg gccatttcag ttctcatgtg tgtactattc tatcattatt gtataacggt 727
231 tttcaaacca gtgggcacac agagaacctc actctgtaat aacaatgagg aatagccacg 787
233 gcgatctcca gcaccaatct ctccatgttt tccacagctc ctccagccaa cccaaatagc 847
235 gcctgctata gtgtagacat cctgcggctt ctagccttgt ccctctctta gtgttcttta 907
237 atcagataac tgcctggaag cctttcattt tacacgccct gaagcagtct tctttgctag 967
239 ttgaattatg tggtgtgttt ttccgtaata agcaaaataa atttaaaaaa atgaaaagtt 1027
242 <210> SEQ ID NO: 14
243 <211> LENGTH: 58
244 <212> TYPE: PRT
245 <213> ORGANISM: Homo sapiens
247 <400> SEQUENCE: 14
248 Met Gln Met Phe Pro Pro Ser Pro Leu Phe Phe Phe Leu Gln Leu Leu
249  1               5                  10                  15
251 Lys Gln Ser Ser Arg Arg Leu Glu His Thr Phe Val Phe Leu Arg Asn
252             20                  25                  30
254 Phe Ser Leu Met Leu Leu Arg Tyr Ile Gly Lys Lys Arg Arg Ala Thr
255         35                  40                  45
257 Arg Phe Trp Asp Pro Arg Arg Gly Thr Pro
258      50                  55
261 <210> SEQ ID NO: 15
262 <211> LENGTH: 111
263 <212> TYPE: PRT
264 <213> ORGANISM: Homo sapiens
266 <400> SEQUENCE: 15
267 Met Lys Thr Asn Asp Thr Tyr Met Lys Phe Ser Trp Leu Thr Val Pro
268  1               5                  10                  15
270 Glu Lys Ser Leu Asp Lys Glu His Arg Cys Ile Val Arg His Glu Asn
271             20                  25                  30
273 Asn Lys Asn Gly Val Asp Gln Glu Ile Ile Phe Pro Pro Ile Lys Thr
274         35                  40                  45
276 Asp Val Ile Thr Met Asp Pro Lys Asp Asn Cys Ser Lys Asp Ala Asn
277      50                  55                  60
279 Asp Thr Leu Leu Leu Gln Leu Thr Asn Thr Ser Ala Tyr Tyr Met Tyr
280  65                  70                  75                  80
282 Leu Leu Leu Leu Leu Lys Ser Val Val Tyr Phe Ala Ile Ile Thr Cys
283                 85                  90                  95
285 Cys Leu Leu Arg Arg Thr Ala Phe Cys Cys Asn Gly Glu Lys Ser
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289 <210> SEQ ID NO: 16
290 <211> LENGTH: 16
291 <212> TYPE: PRT
292 <213> ORGANISM: Homo sapiens
294 <220> FEATURE:
295 <223> OTHER INFORMATION: Partial amino acid sequence of TARP (residues
296       42-57)
298 <400> SEQUENCE: 16
299 Gly Lys Lys Arg Arg Ala Thr Arg Phe Trp Asp Pro Arg Arg Gly Thr
300   1               5                  10                  15
303 <210> SEQ ID NO: 17
304 <211> LENGTH: 16
305 <212> TYPE: PRT
306 <213> ORGANISM: Dictyostelium discoideum
308 <220> FEATURE:
309 <223> OTHER INFORMATION: Partial amino acid sequence of Tup1 (dTup,
310       residues 521-536)
312 <400> SEQUENCE: 17
313 Gly Ser Lys Asp Arg Ser Val Gln Phe Trp Asp Pro Arg Asn Gly Thr
314   1               5                  10                  15
317 <210> SEQ ID NO: 18
318 <211> LENGTH: 16
319 <212> TYPE: PRT
320 <213> ORGANISM: Saccharomyces cerevisiae
322 <220> FEATURE:
323 <223> OTHER INFORMATION: Partial amino acid sequence of Tup1 (yTup1,
324       residues 626-660)
326 <400> SEQUENCE: 18
327 Gly Ser Lys Asp Arg Gly Val Leu Phe Trp Asp Lys Lys Ser Gly Asn
328   1               5                  10                  15
331 <210> SEQ ID NO: 19
332 <211> LENGTH: 41
333 <212> TYPE: DNA
334 <213> ORGANISM: Artificial Sequence
336 <220> FEATURE:
337 <223> OTHER INFORMATION: Description of Artificial Sequence:PCR primer
339 <400> SEQUENCE: 19
340 ttacagataa acaacttgat acagatgttt cccccaagcc c                    41
343 <210> SEQ ID NO: 20
344 <211> LENGTH: 39
345 <212> TYPE: DNA
346 <213> ORGANISM: Artificial Sequence
348 <220> FEATURE:
349 <223> OTHER INFORMATION: Description of Artificial Sequence:PCR primer
351 <400> SEQUENCE: 20
352 gggcttgggg gaaacatctg tatcaagttg tttatctgt                       39
355 <210> SEQ ID NO: 21
356 <211> LENGTH: 36
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VERIFICATION SUMMARY DATE: 11/21/2005
PATENT APPLICATION: **US/10/031,158B** TIME: 09:58:30

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